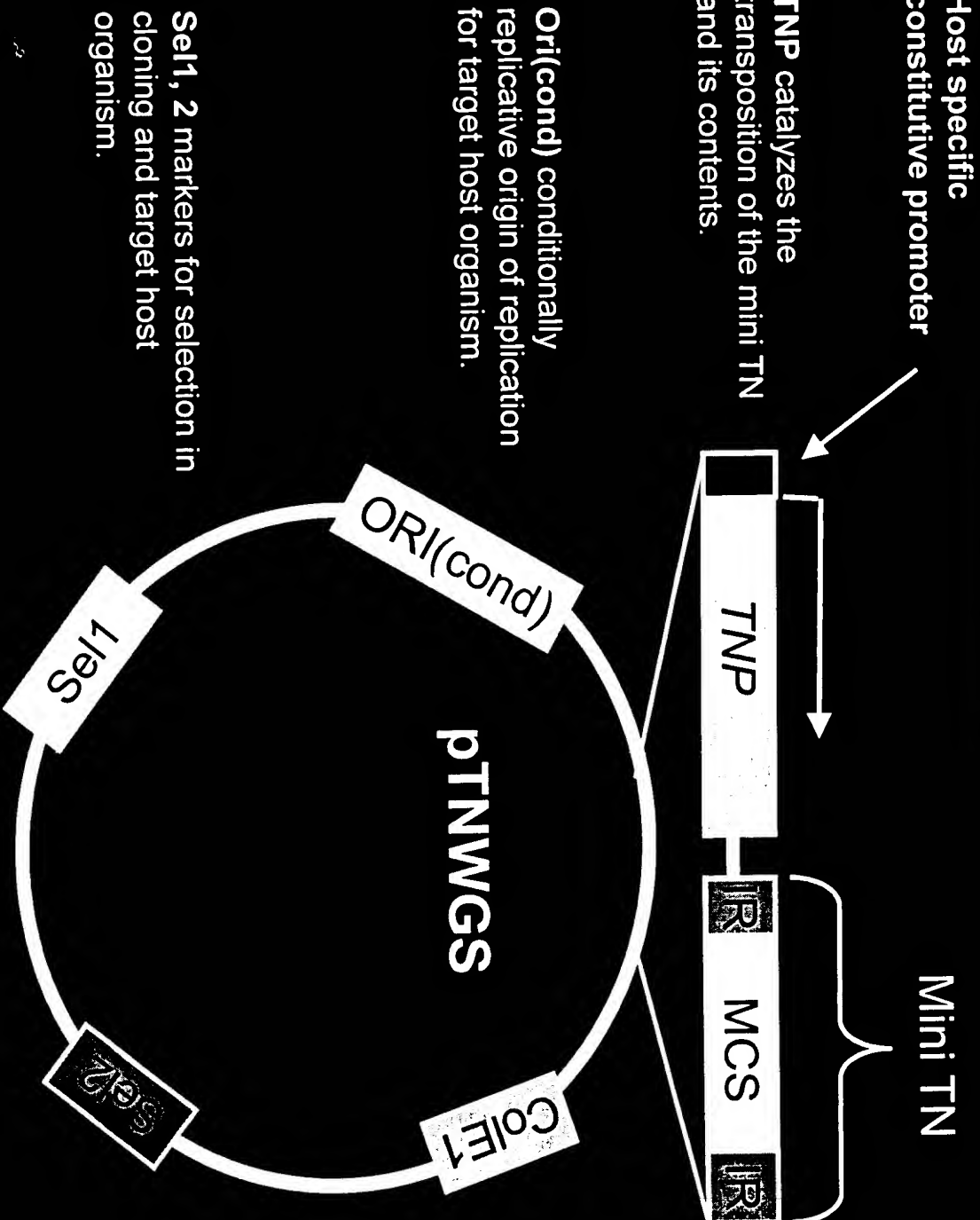


pTNMAX (general vector)



Mini-TN inverted repeats from a transposon active in the target host organism.

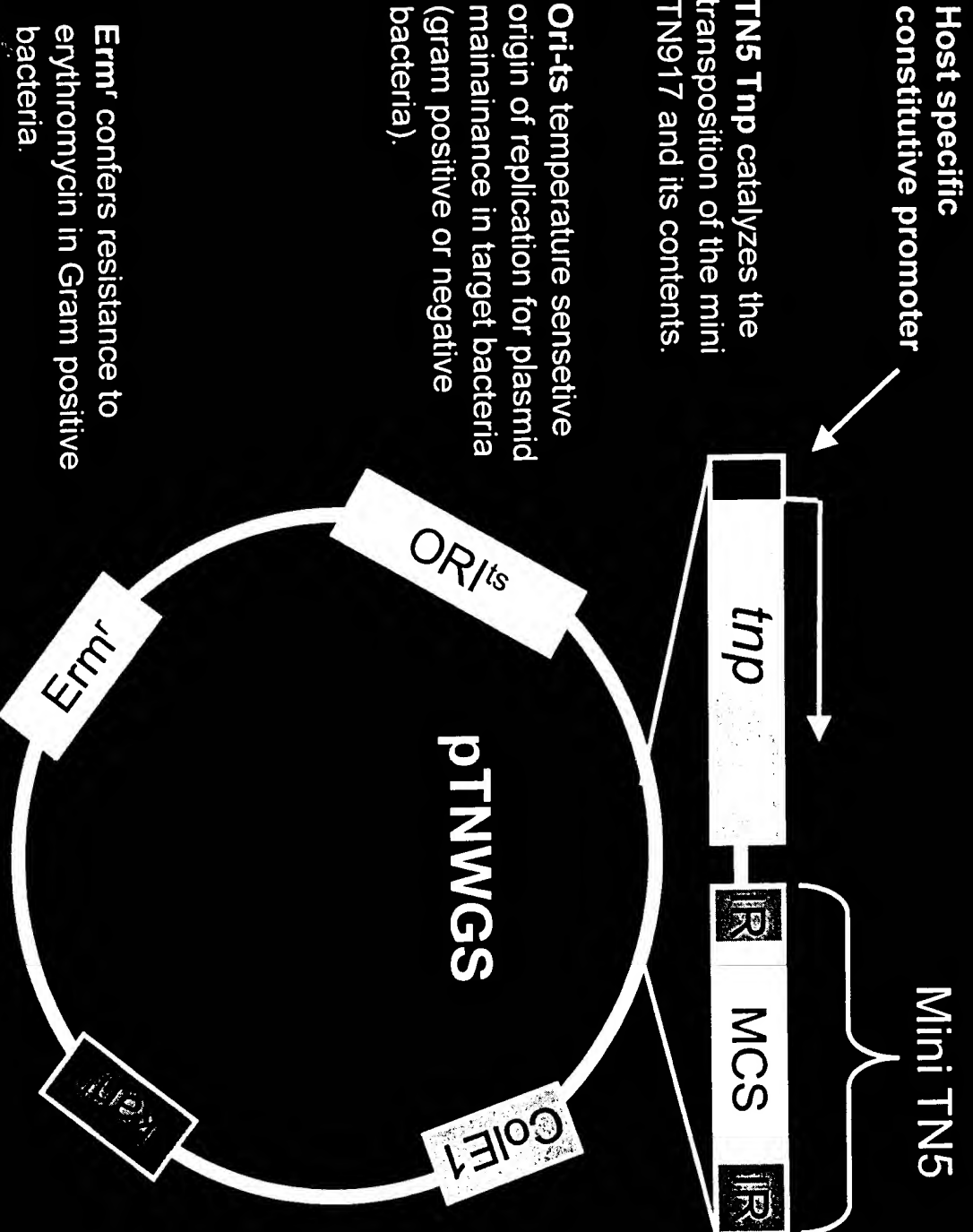
MCS multiple cloning site for the introduction of gene libraries into the mini ISS1 transposon.

ORI – origin of replication for the desired cloning host organism.

Se1, 2 markers for selection in cloning and target host organism.

Figure 1 A

pWGS:5



Mini TN5 TN5 inverted repeats flanking a multiple cloning site into which gene libraries can be cloned.

ColE1 origin of replication for plasmid maintenance in *E. coli*.

Kan^r confers resistance to kanamycin to *E. coli*

Figure 1B

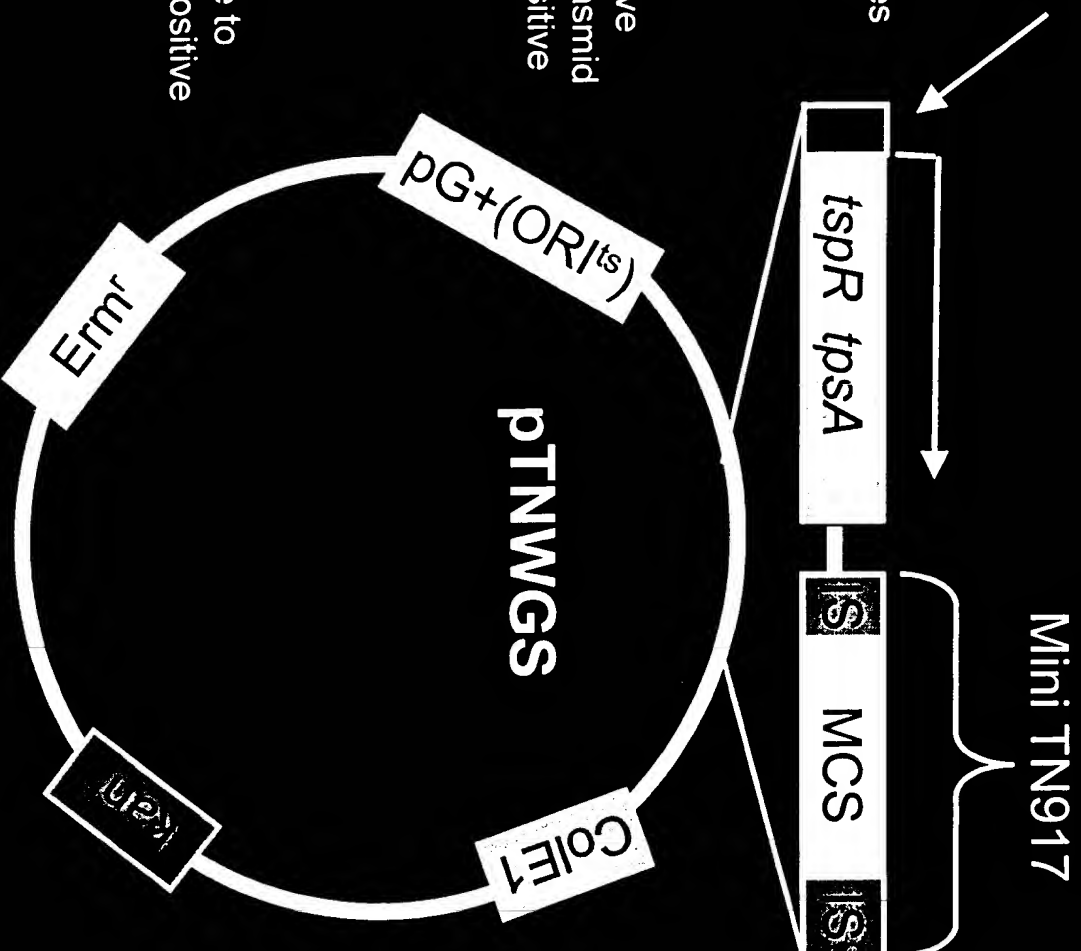
pWGS:917

Host specific promoter – nisaA
promoter for lactic acid bacteria.

917 TspR TspA catalyzes a transposition of the mini 917 and its contents (transposase/resolvase)

PG+ temperature sensitive origin of replication for plasmid maintenance in Gram positive bacteria.

Ernr confers resistance to erythromycin in Gram positive bacteria.



MCS multiple cloning site for the introduction of gene libraries into the mini 917 transposon.

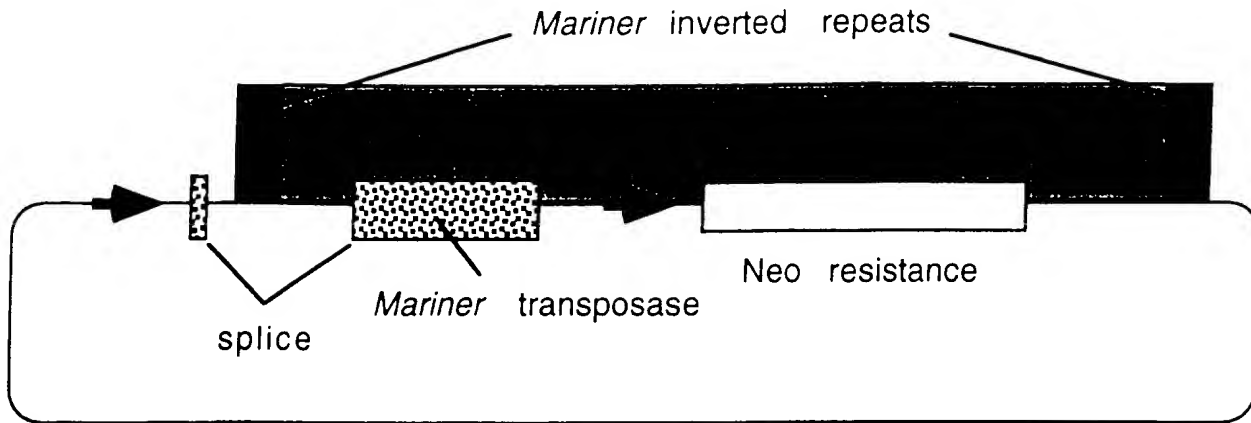
COIE1 origin of replication for plasmid maintenance in *E. coli*.

Kanr confers resistance to kanamycin to *E. coli*.

Figure 1C

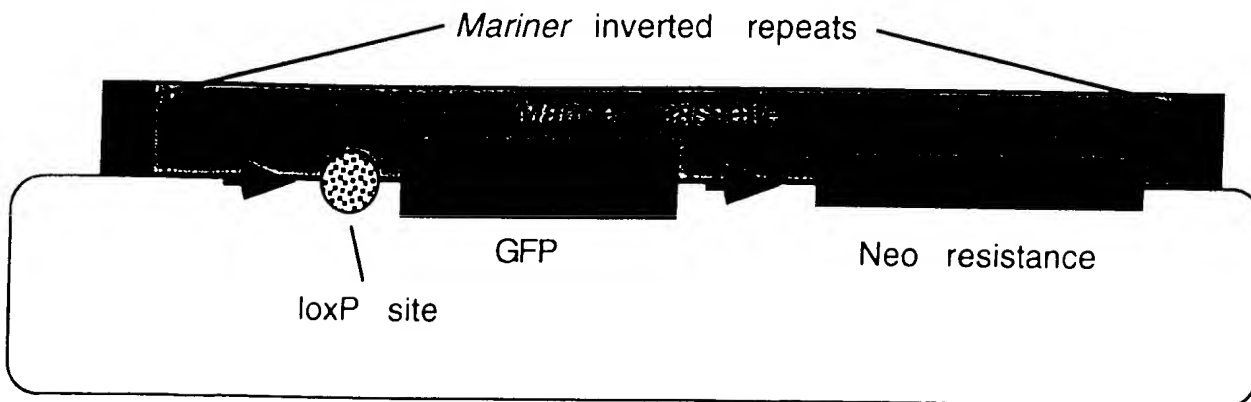
A

Efficient integration into mammalian cells using evolved *Mariner* transposons



B

Mariner transposon for inserting loxP sites at loci with desirable expression properties



Methodology for Isolating Hosts with improved Phenotypes by Whole Genome Shuffling (WGS)

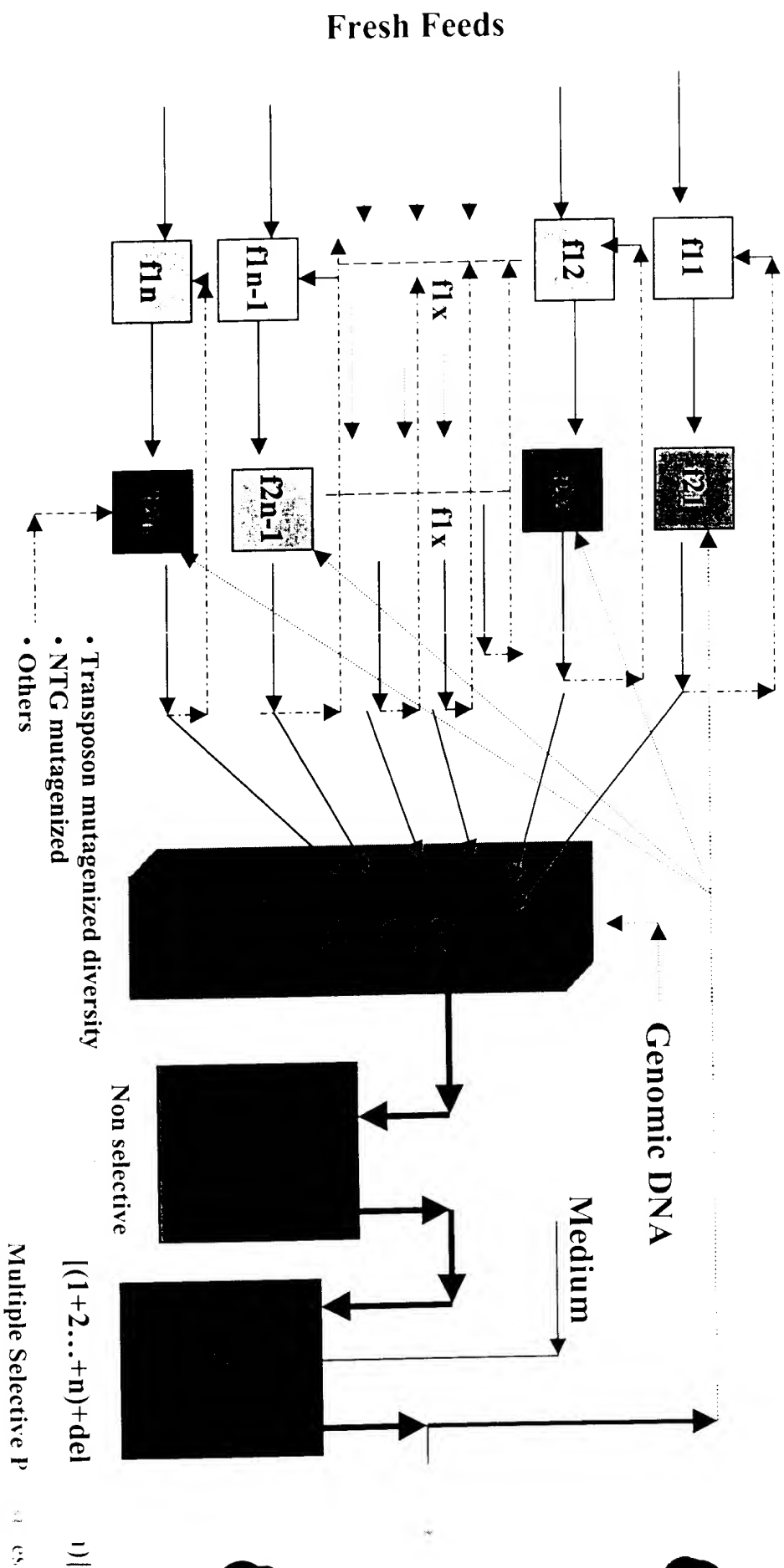


Figure 3

Shuffling of Genomes *In Vitro*: Formation of transposomes

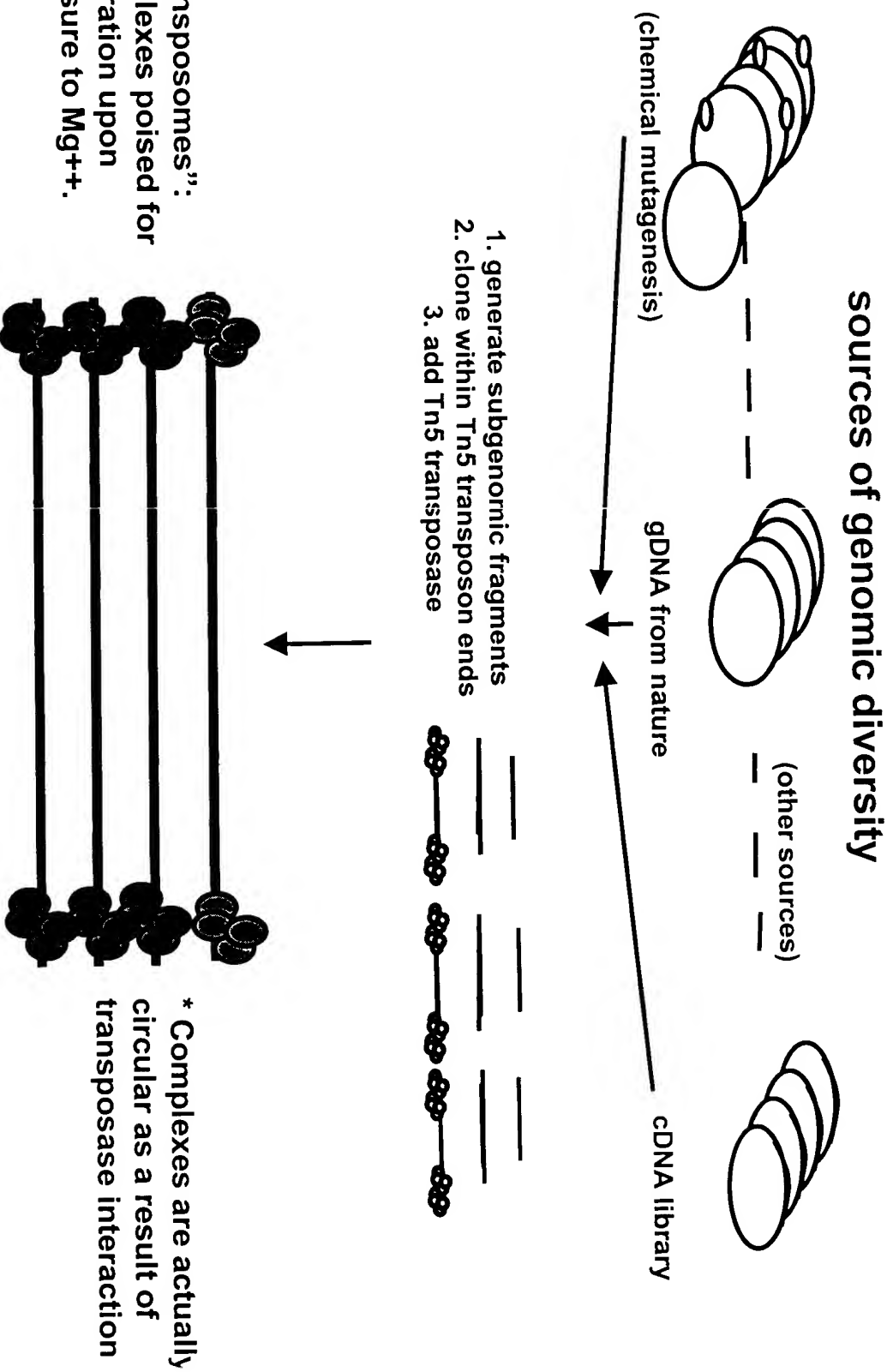


Figure 4A

Shuffling of Genomes *In Vitro*: Breeding multiple donor genomes with a single acceptor genome

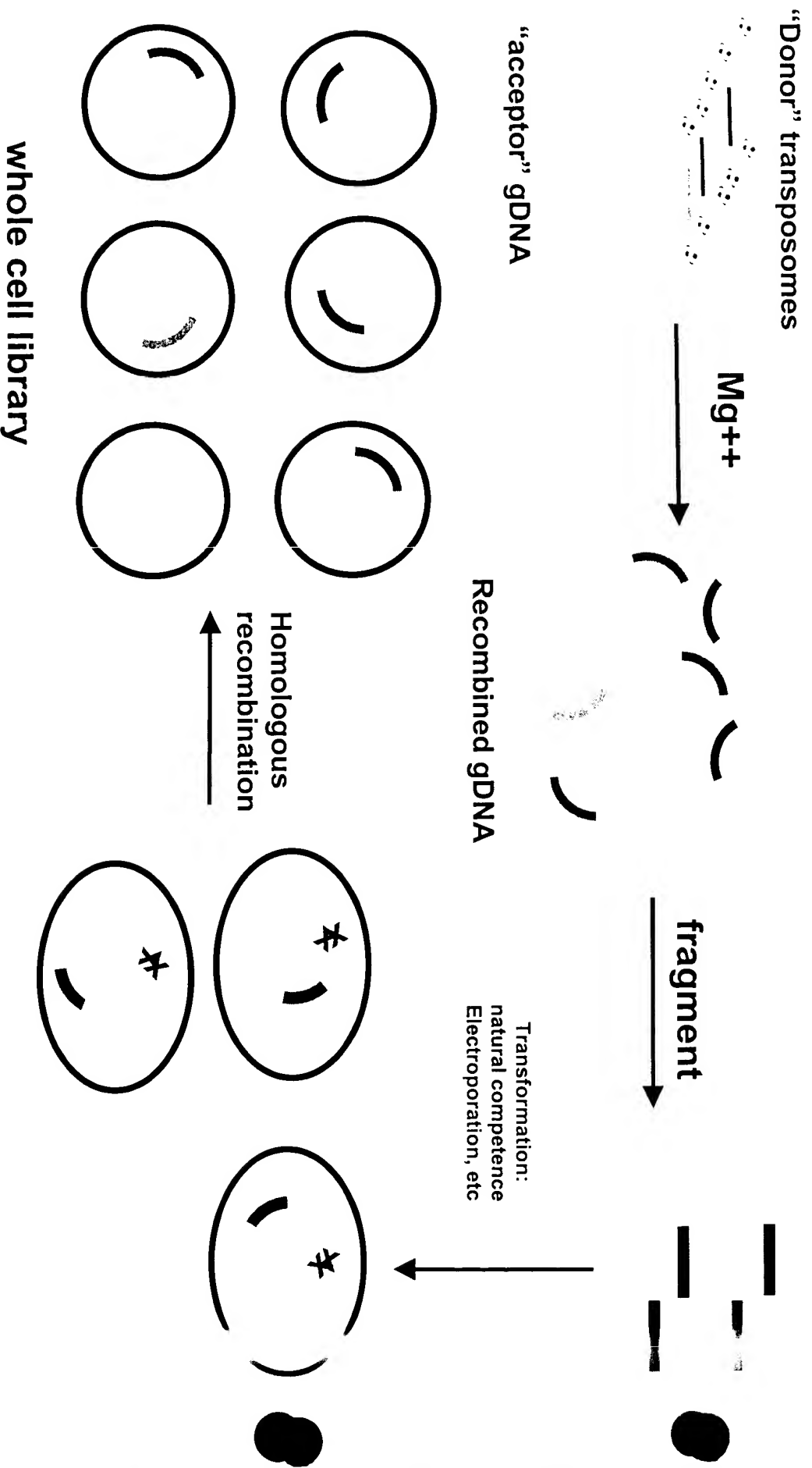


Figure 4B

Shuffling of Genomes *In Vitro*: Breeding multiple donor genomes with multiple acceptor genomes

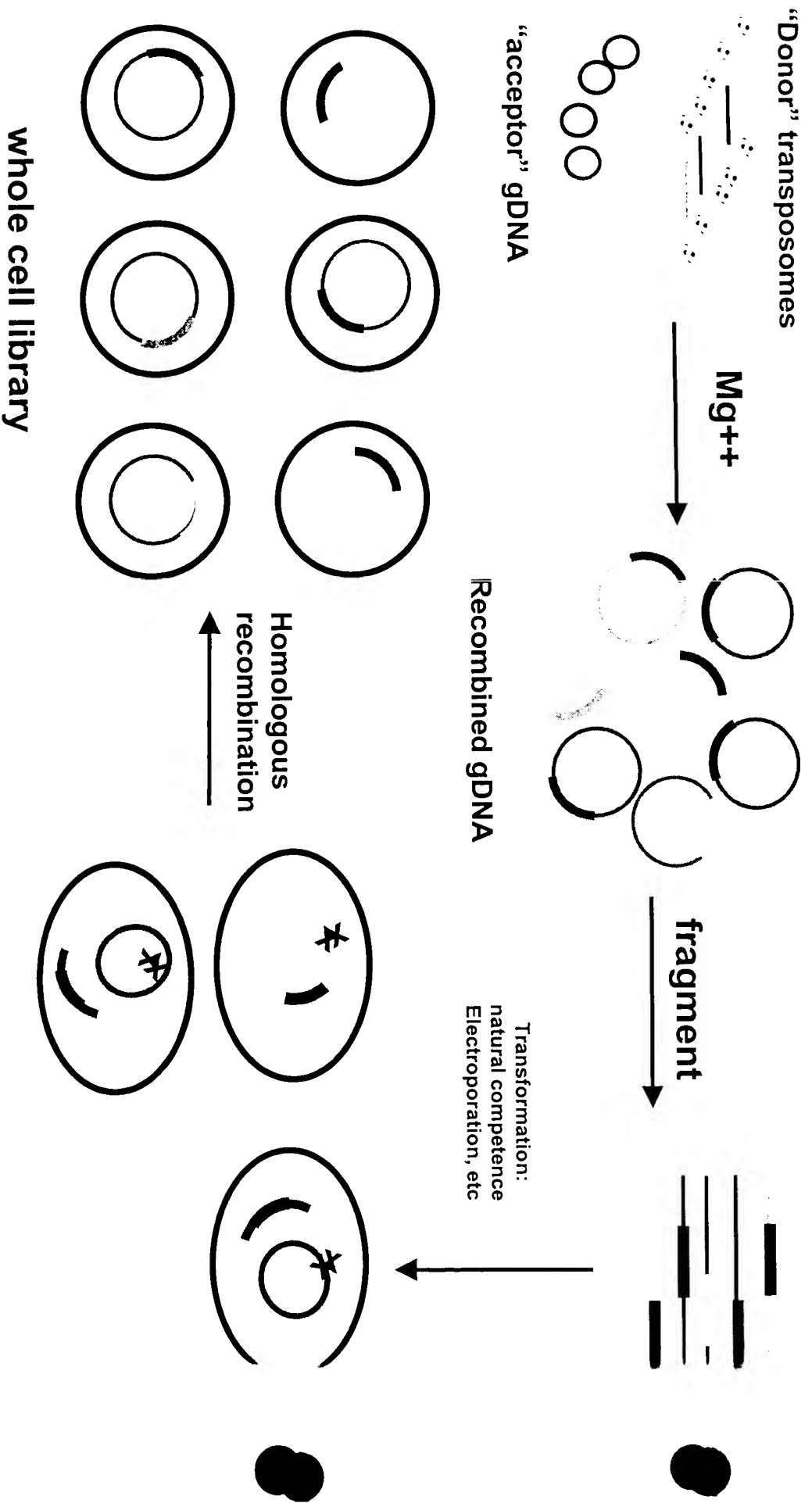


Figure 4C

Shuffling of Genomes *In Vitro*: Split pool recursive *in vitro* recombination of multiple genomes

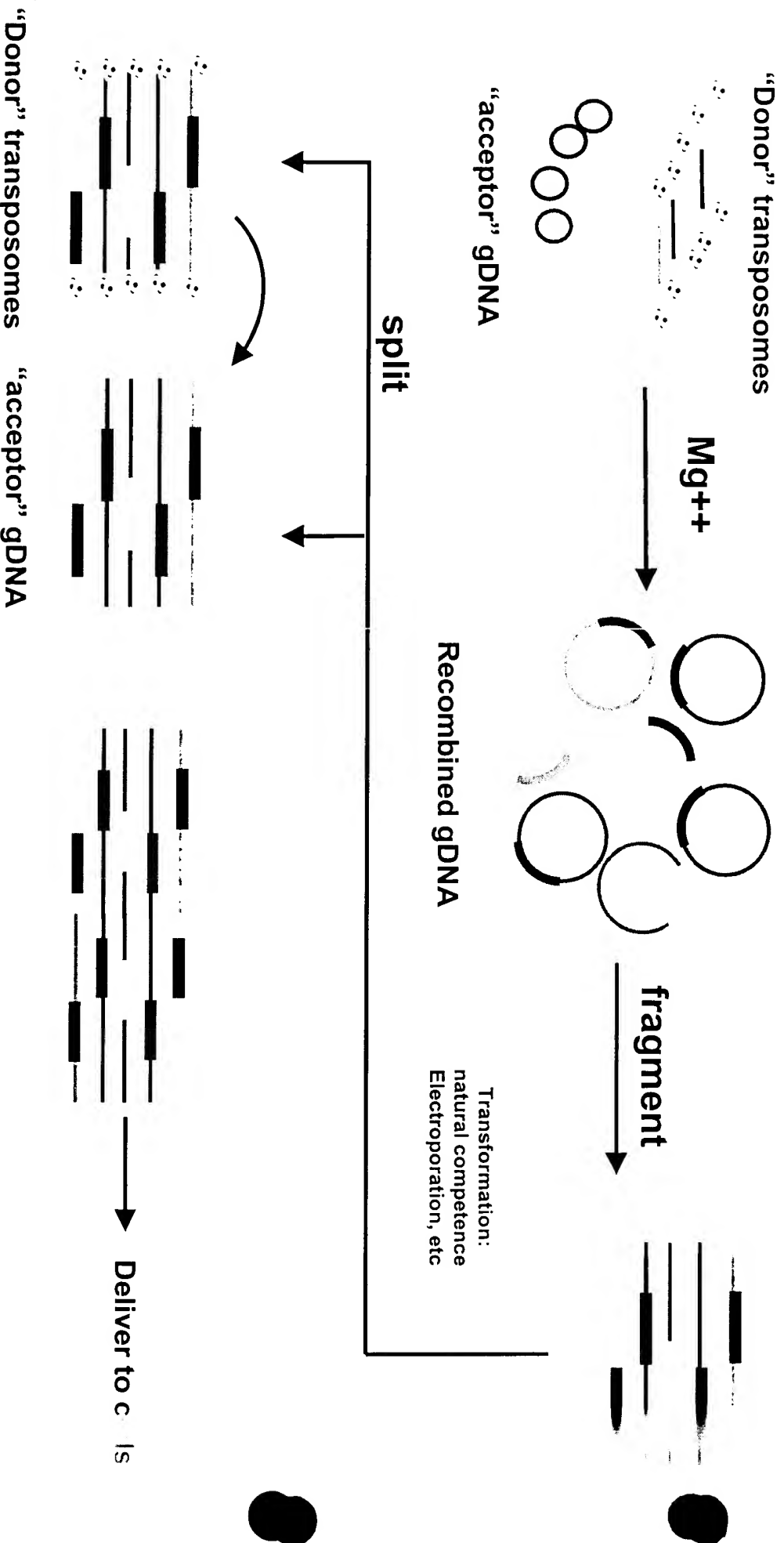


Figure 4D